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François Rousseau

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Research Report

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A Non-Local Approach for Image Super-Resolution using Intermodality Priors

F. Rousseau



A Non-local Approach for Image Super-Resolution using Intermodality Priors

F. Rousseau and the Alzheimer’s Disease Neuroimaging Initiative *
LSIIT, UMR 7005 CNRS - Université de Strasbourg, France

Abstract

Image enhancement is of great importance in medical imaging where image resolution remains a crucial point in many image analysis algorithms. In this paper, we investigate brain hallucination [22], or generating a high resolution brain image from an input low-resolution image, with the help of another high resolution brain image. We propose an approach for image super-resolution by using anatomical intermodality priors from a reference image. Contrary to interpolation techniques, in order to be able to recover fine details in images, the reconstruction process is based on a physical model of image acquisition. Another contribution to this inverse problem is a new regularization approach that uses an example-based framework integrating non-local similarity constraints to handle in a better way repetitive structures and texture. The effectiveness of our approach is demonstrated by experiments on realistic Brainweb Magnetic Resonance images and on clinical images from ADNI, generating automatically high-quality brain images from low-resolution input.

1 Introduction

Image processing algorithm performances are often limited by the image resolution. Resolution is for instance a key point in brain segmentation in Magnetic Resonance (MR) imaging for which partial volume effect is a limiting factor for fine image analysis. Then, it clearly appears that improving image resolution is still one of the main challenges in medical image processing. In medical imaging, a so-called low resolution (LR) 3D image is usually a stack of 2D thick slices. As a result, 3D data are generally not isotropic. Figure 1 shows for instance a high resolution (HR) T1-weighted image and a LR T2-weighted image of the same patient (in this case, these images come from the Brainweb database [6]). In many cases, it may be necessary to put HR and LR images into a same coordinate system. In medical imaging, this is usually done by applying interpolation techniques [15]. Image interpolation is a very common image processing technique in medical imaging pipelines and it may have a strong impact on other processing steps such as segmentation or registration. Interpolation methods can be divided into two groups: scene-based and object-based methods [12]. Scene-based approaches use only image intensities to determine the interpolated intensity (for instance: nearest neighbor, linear interpolation, spline-based interpolation). Such scene-based methods produce perceptually unsatisfactory results with blurred edges and textures. Many edge-preserving interpolation techniques have been reported to handle this problem. However, these techniques rely on accurate edge information that is not obtainable from coarse data. In order to guide the interpolation process, object-based methods make use of additional information extracted from images. As an example, non-rigid registration techniques have been used to drive interpolation methods between adjacent slices [9], [19]. However, scene-based and object-based techniques do not take advantage of a model of the imaging process.

Super-resolution (SR) is another model-based technique which relies on modeling the imaging processes and using regularization methods describing *a priori* constraints. The principle of this approach is usually combining LR images to produce an image that has a higher spatial resolution than the original images [3]. In medical imaging, several SR methods have been proposed to combine LR images to reconstruct one HR image [18, 5, 21] (usually by modifying the acquisition protocol). SR is a large research field encompassing many applications. The work we present in this paper is related to the single-frame SR framework [23], meaning that only one LR image is used to generate an HR image. More specifically, we focus on studies involving MR imaging for which an anatomical HR image and several other LR images are acquired to keep acquisition

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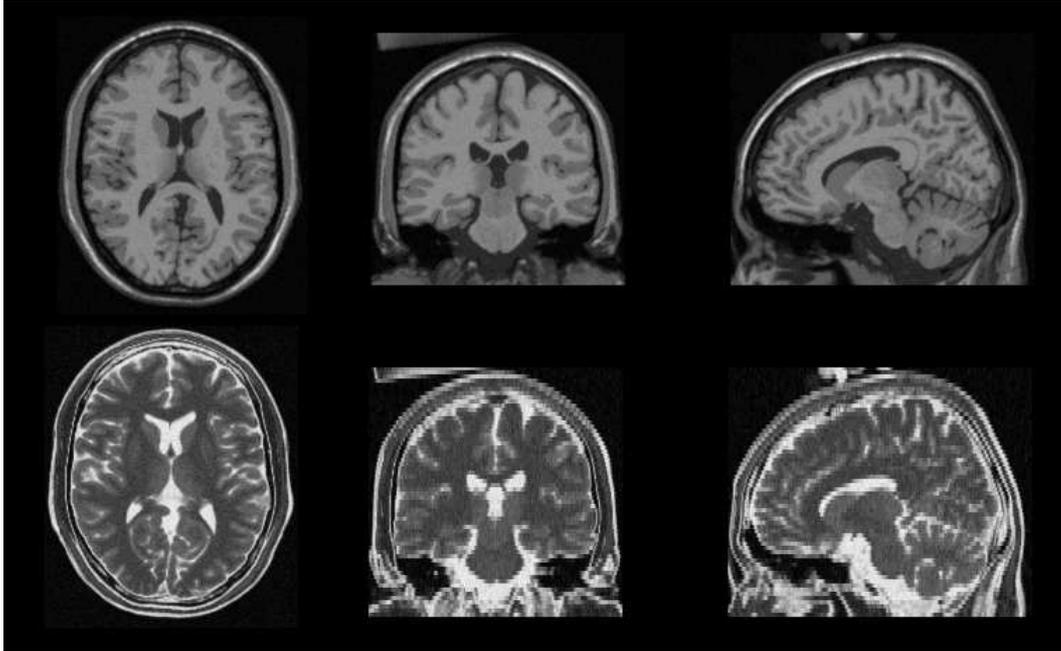


Figure 1: Example of MR Brainweb data [6]. First row: high resolution T1-weighted image ($1 \times 1 \text{mm}^2$ in-plane resolution, 1mm slice thickness); second row: low resolution T2-weighted image ($1 \times 1 \text{mm}^2$ in-plane resolution, 3mm slice thickness). The purpose of the proposed approach is to reconstruct a high-resolution T2-weighted image using information from the T1-weighted image.

time at an acceptable level for the patient (see Figure 1). This is the case for routinely performed clinical MR acquisitions such as follow-up of neurodegenerative diseases, brain tumor evolution or diffusion tensor imaging. Typically, one isotropic HR T1-weighted image and several anisotropic LR images (such as T2-weighted, FLAIR or proton density images) are acquired. One can note that this issue of relative difference in image resolution may also occur in other clinical settings. For instance, patient follow-up may require an analysis of scans taken at different magnetic field strength, potentially at different clinical sites.

In such context, we propose a new approach for image SR by using information from an HR MR image to drive the image reconstruction of the LR MR image. The general framework developed in this paper can be applied to other domains where image resolution is an important issue, such as remote sensing. This paper is built on previously published work [22]. Many experiments have been added in order to provide further insight to the proposed approach. In Section 2, we present the image SR problem using a model-based framework and some recently proposed example-based approaches. Section 3 details our non-local approach for image SR using an HR reference MR image. In Section 4, results obtained on realistic brainweb images are presented and discussed.

2 Image super-resolution

In this section, we present the model-based framework for image SR which leads to an ill-posed inverse problem. Then, recently proposed regularization approaches are described.

2.1 Model-based framework

Contrary to interpolation approaches, model-based approaches use a generic observation model such as:

$$\mathbf{y} = DBW\mathbf{x} + \mathbf{n} \quad (1)$$

where \mathbf{y} denotes the LR image, \mathbf{x} is the HR image, \mathbf{n} represents observation noise, D is the sub-sampling matrix, B a blur matrix, W is the geometric transformation.

The three operators can be combined into a single matrix H : $H = DBW$. The matrix H thus incorporates motion compensation, degradation effects, and sub-sampling for the LR image \mathbf{y} . In this paper, we assume that the degradation operator H and the noise characteristics are known and we focus only on the reconstruction step.

Based on this observation model, the SR image can be estimated by minimizing a least-square cost function such as:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \|\mathbf{y} - H\mathbf{x}\|^2. \quad (2)$$

For such an inverse problem, some form of regularization plays a crucial role and must be included in the cost function to stabilize the problem or constrain the space of solutions [13]. Thus, the HR image is computed by considering the following equation:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \mathcal{L}(\mathbf{x}, \mathbf{y}, H) + \lambda \mathcal{R}(\mathbf{x}). \quad (3)$$

$\mathcal{L}(\mathbf{x}, \mathbf{y}, H)$ is a data fidelity term related to the physical model that penalizes inconsistency between the estimated HR image \mathbf{x} and the observed LR image \mathbf{y} . This term depends on the observation model used. In our case, the data fidelity term is simply defined as following:

$$\mathcal{L}(\mathbf{x}, \mathbf{y}, H) = \|\mathbf{y} - H\mathbf{x}\|^2. \quad (4)$$

$\mathcal{R}(\mathbf{x})$ is a regularization term. A common approach for regularization is to take explicitly into account the image geometry and to introduce a global weight λ that balances the contribution of prior smoothness terms and a fidelity term.

2.2 Image regularization

Examples of popular pixel-based regularizers are:

- Tikhonov regularization: $\mathcal{R}(\mathbf{x}) = \sum_{p=0}^P \int_{\Omega} c_p |\mathbf{x}^{(p)}|^2$ where c_p are positive coefficients, Ω is the domain of integration (all the voxels) and $\mathbf{x}^{(p)}$ is p^{th} -order derivative of \mathbf{x} .
- Variational approach: $\mathcal{R}(\mathbf{x}) = \sum_{p=0}^P \int_{\Omega} c_p |\mathbf{x}^{(p)}|^q$ where q is different from 2 ($q = 2$ is the Tikhonov regularization).
- Markov random field *a priori* image model: $\mathcal{R}(\mathbf{x}) = \alpha \sum_s \phi(\mathbf{x}_s) + \beta \sum_{r,s} \phi(\mathbf{x}_r - \mathbf{x}_s)$ where ϕ can be a L_2L_1 function for instance.

However, the use of such a regularization term assumes a specific image model: for instance, variational approaches can be based on the assumption that images are made of smooth regions separated by sharp edges. Such prior regularization approach introduces then strong constraints into the reconstruction process.

Explicit models for the many regularities and geometries seen in local patterns are needed to develop better image reconstruction algorithms. In contrast to the pixel-based regularization approach discussed above, example-based methods consists of modeling non-local pairwise interactions from training data or a library of image patches. The principle of example-based SR methods is to add a similarity constraint between the voxels of HR image and the nearest examples present in the learning database [2],[8]. Such an approach suggests that the reconstructed image should locally look like examples existing in the learning database. The regularization term can then be defined in a general way as follows:

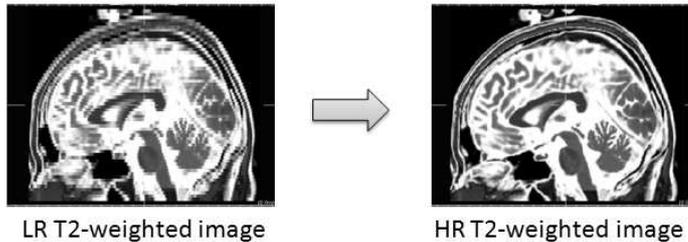
$$\mathcal{R}(\mathbf{x}, \mathcal{E}) = \sum_{\mathbf{v}, \mathbf{k} \in \Omega(\mathbf{v})} w_{\mathbf{v}, \mathbf{k}} \|f(\mathbf{x}(\mathbf{v})) - \mathcal{E}_{\mathbf{v}, \mathbf{k}}\|^2 \quad (5)$$

where $f(\mathbf{x}(\mathbf{v}))$ is an operator on the HR image \mathbf{x} at the voxel \mathbf{v} , $\Omega(\mathbf{v})$ is a neighborhood of \mathbf{v} , \mathcal{E} is the learning database, $\mathcal{E}_{\mathbf{v}, \mathbf{k}}$ is a element of \mathcal{E} related to \mathbf{v} and $w_{\mathbf{v}, \mathbf{k}}$ is a local weight. It is important to note that in example-based methods the regularization term \mathcal{R} depends on the learning database \mathcal{E} . For instance (see [2]), $f(\mathbf{x}(\mathbf{v}))$ can stand for the gradient value of the SR image \mathbf{x} at the voxel \mathbf{v} and $\mathcal{E}_{\mathbf{v}, \mathbf{k}}$ would be the gradient value of the best matching pixel with respect to \mathbf{v} in the learning database.

Although the possibility to introduce new image priors makes the example-based approach very attractive, the key point (which can be a major drawback depending on the application) is the need of a relevant learning database.

In this work, we proposed to take advantage of the existence of an HR reference image in the MR imaging context and to investigate the use of a patch-based approach without any learning database by assuming that there exist related patterns in the LR anisotropic image and an HR isotropic image of the same patient. From the point of view of example-based techniques, the HR isotropic image can be seen as a specific learning image database since information present in the HR image is used to constrain the ill-posed reconstruction problem.

Single image super-resolution



Proposed super-resolution



Figure 2: Comparison of single image SR framework and the proposed SR approach. Single image SR techniques aim at estimating an HR image using only one LR image as input. In the proposed strategy, the SR estimation is done using jointly the LR image and a reference HR image.

3 Proposed Approach

3.1 Overview

The main idea of this work is to reconstruct an HR image using one LR image and intermodality priors from another HR image. In this context, we propose to use a non-local patch-based approach to define the regularization term in order to take into account complex spatial interactions within images. Moreover, in contrast to example-based approaches for image modeling, the proposed method is unsupervised and thus uses no image patch learning database and no computationally intensive training algorithms.

The three key points of the proposed approach are:

- a non-local regularization functional which can handle repetitive patterns in the image domain,
- the assumption that an HR reference image \mathcal{E}_x of a patient may contain relevant examples which should be used to reconstruct an HR image \mathbf{x} from a LR image \mathbf{y} of the same patient,
- no use of external learning database.

The principle of non-local techniques is briefly described in Section 3.2. Section 3.3 presents how the patterns of the reference image \mathcal{E}_x are used to drive the reconstruction process of \mathbf{x} . An overview of the algorithm is provided in Section 3.4.

3.2 A non-local approach

Recently, Buades *et al.* [4] have proposed a very efficient denoising algorithm relying on a non-local framework. Since then, this non-local strategy has been studied and applied in several image processing applications such as non-local regularization functionals in the context of inverse problems [22, 17, 14, 11, 20].

Let us consider a weighted graph w that links together voxels \mathbf{v} , \mathbf{k} over the image domain with a weight $w(\mathbf{v}, \mathbf{k})$. This weighted graph w is a representation of non-local interactions between image elements. In [4], the graph w is used for denoising purpose using a non-local neighborhood averaging strategy (called non-local

means (NLM)):

$$\forall \mathbf{v}, d_{NLM}(\mathbf{x}(\mathbf{v})) = \sum_{\mathbf{k} \in \Omega(\mathbf{v})} w(\mathbf{v}, \mathbf{k}) \mathbf{x}(\mathbf{k}) \quad (6)$$

where $d_{NLM}(\mathbf{x}(\mathbf{v}))$ is a denoised version of $\mathbf{x}(\mathbf{v})$. The computation of the weighted graph is described in the Appendix.

Buades *et al.* have shown that, for 2D natural images, the NLM filter outperforms state-of-the-art denoising methods such as the Rudin-Osher-Fatemi Total Variation minimization scheme or the Perona-Malik Anisotropic diffusion. The NLM method tries to take advantage of the high degree of redundancy of any natural image and appears to be an unsupervised example-based denoising method.

In [22, 17], a non-local regularization functional for inverse problems has been proposed relying on a NLM denoised version of the reconstructed image:

$$\mathcal{R}(\mathbf{x}) = \sum_{\mathbf{v}} \|\mathbf{x}(\mathbf{v}) - d_{NLM}(\mathbf{x}(\mathbf{v}))\|^2. \quad (7)$$

Such a regularization functional has the advantage over popular variational regularization functionals to handle in a better way repetitive structures and texture without incorporating strong priors on the reconstructed image.

3.3 A non-local regularization functional using intermodality priors

We propose to introduce into the regularization term a similarity constraint between local patterns of the reconstructed HR image \mathbf{x} and the HR reference image $\mathcal{E}_{\mathbf{x}}$.

Let $w_{\mathcal{E}_{\mathbf{x}}}$ the weighted graph estimated on the HR reference image $\mathcal{E}_{\mathbf{x}}$ and $w_{\mathbf{x}}$ the weighted graph estimated on the HR reconstructed image \mathbf{x} .

A naive approach to use intermodality priors for SR would be to directly compute the weighted graph $w_{\mathcal{E}_{\mathbf{x}}}$ and to estimate the SR image using the regularization functional defined in Equation 7. In this case, the denoised version of the reconstructed HR image \mathbf{x} would be computed using the weighted graph $w = w_{\mathcal{E}_{\mathbf{x}}}$ in Equation 6:

$$\forall \mathbf{v}, d_{NLM}(\mathbf{x}(\mathbf{v}), \mathcal{E}_{\mathbf{x}}) = \sum_{\mathbf{k} \in \Omega(\mathbf{v})} w_{\mathcal{E}_{\mathbf{x}}}(\mathbf{v}, \mathbf{k}) \mathbf{x}(\mathbf{k}) \quad (8)$$

This simple strategy relies on the assumption that local patterns in the HR reference image $\mathcal{E}_{\mathbf{x}}$ could be directly used as examples to regularize the reconstructed HR image \mathbf{x} ¹. However, there are some cases where this assumption may not hold. In the context of MR imaging, images $\mathcal{E}_{\mathbf{x}}$ and \mathbf{y} are not acquired with the same MR sequence. Multimodal MR data do not reveal the same tissue specificity; for instance, Multiple Sclerosis (MS) lesions are clearly visible in T2-weighted images but not in T1-weighted images (see Figure 3). In this case, the HR T1-weighted image may not be a relevant candidate to guide the reconstruction process. In order to handle the case of possible *outliers*, we propose to modify the regularization term by locally analyzing the correlation between the two graphs $w_{\mathbf{x}}$ and $w_{\mathcal{E}_{\mathbf{x}}}$.

For each voxel \mathbf{v} , we define a scalar α as the correlation between the two weighted graphs $w_{\mathbf{x}}$ and $w_{\mathcal{E}_{\mathbf{x}}}$. To handle outliers (such as MS lesions), we propose an adaptive scheme by modifying the way to compute the denoised version of the HR reconstructed image \mathbf{x} :

$$d_{NLM}(\mathbf{x}(\mathbf{v}), \mathcal{E}_{\mathbf{x}}) = \alpha(\mathbf{v}) \sum_{\mathbf{k} \in \Omega(\mathbf{v})} w_{\mathcal{E}_{\mathbf{x}}}(\mathbf{v}, \mathbf{k}) \mathbf{x}(\mathbf{k}) + (1 - \alpha(\mathbf{v})) \sum_{\mathbf{k} \in \Omega(\mathbf{v})} w_{\mathbf{x}}(\mathbf{v}, \mathbf{k}) \mathbf{x}(\mathbf{k}). \quad (9)$$

$d_{NLM}(\mathbf{x}(\mathbf{v}), \mathcal{E}_{\mathbf{x}})$ is now a weighted average of two denoised versions of $\mathbf{x}(\mathbf{v})$. The first term is the denoised version of $\mathbf{x}(\mathbf{v})$ computed by using the HR reference image $\mathcal{E}_{\mathbf{x}}$ and the second term is a denoised version of $\mathbf{x}(\mathbf{v})$ obtained with the current estimate of the reconstructed HR image \mathbf{x} . If the two weighted graphs are correlated, the HR reference image $\mathcal{E}_{\mathbf{x}}$ is likely to be a relevant candidate to guide the reconstruction process and thus, α is close to 1. If the weighted graphs are uncorrelated, the presence of outliers is detected and α is close to 0. Defining $d_{NLM}(\mathbf{x}(\mathbf{v}), \mathcal{E}_{\mathbf{x}})$ in such way allows us to choose the best examples to regularize the reconstructed HR image \mathbf{x} .

We define then a regularization functional $\mathcal{R}(\mathbf{x}, \mathcal{E}_{\mathbf{x}})$ taking advantage of intermodality priors as follows:

$$\mathcal{R}(\mathbf{x}, \mathcal{E}_{\mathbf{x}}) = \sum_{\mathbf{v}} \|\mathbf{x}(\mathbf{v}) - d_{NLM}(\mathbf{x}(\mathbf{v}), \mathcal{E}_{\mathbf{x}})\|^2 \quad (10)$$

where $d_{NLM}(\mathbf{x}(\mathbf{v}), \mathcal{E}_{\mathbf{x}})$ is a denoised version of $\mathbf{x}(\mathbf{v})$ using the HR reference image $\mathcal{E}_{\mathbf{x}}$ (see Equation 9).

¹It is important to note that in example-based SR methods, the elements of the learning database used for reconstruction are image intensities. In the proposed approach, the elements of the learning database are weighted graphs, which describe non-local interactions between image voxels.

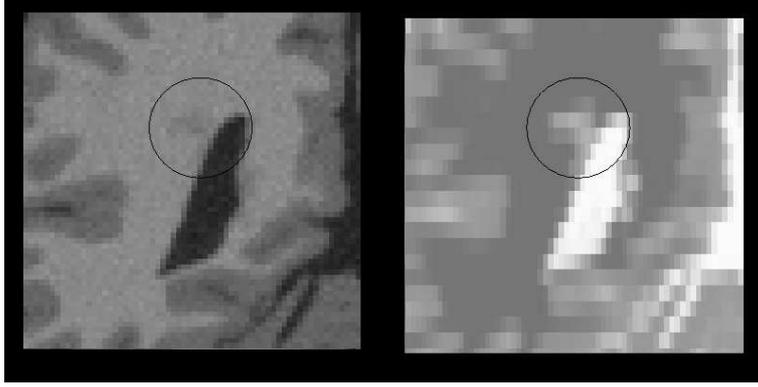


Figure 3: Illustration of possible outliers. The MS lesion (in the circle) is less visible in the HR T1-weighted image than in the LR T2-weighted image. In this case, the assumption that local patterns in the HR image \mathcal{E}_x could be used as examples to regularize the reconstructed HR image \mathbf{x} does not hold.

3.4 Overview of the algorithm

Require: LR image \mathbf{y} , starting estimate of HR image \mathbf{x} , HR reference image \mathcal{E}_x

- 1: Compute the weighted graph $w_{\mathcal{E}_x}$
 - 2: **repeat**
 - 3: Compute the weighted graph w_x
 - 4: Update α
 - 5: Estimate $d_{NLM}(\mathbf{x}, \mathcal{E}_x)$
 - 6: Update \mathbf{x} by minimizing $\mathcal{L}(\mathbf{x}, \mathbf{y}, H) + \lambda \mathcal{R}(\mathbf{x}, \mathcal{E}_x)$
 - 7: **until** convergence
-

4 Results

To explore the ability to reconstruct high resolution images of realistic typical anatomical brain structures, we applied the algorithm on MRI images of Brainweb [6] and real images (ADNI data).

4.1 Brainweb images

Brainweb is a simulated brain database which is often used as a gold standard for the analysis of in vivo MR data. Using the HR image provided by Brainweb, we have generated LR images with an in-plane resolution of 1mm x 1mm and a slice thickness of 3mm, using the observation model described by Equation 1. As in the Brainweb simulator, the point spread function is modeled with a boxcar function. Intensities of HR Brainweb images have been normalized between 0 and 255.

The database contains simulated brain MRI data based on two anatomical models: normal and multiple sclerosis (MS lesions have been extracted from real MRI data). We also have investigated the influence of Gaussian noise (in this case, both LR images and HR reference images are corrupted with Gaussian noise). The proposed methodology has been implemented on 4 cases:

1. normal anatomical model without noise,
2. multiple sclerosis anatomical model without noise,
3. normal anatomical model with Gaussian noise ($\sigma = 10$),
4. multiple sclerosis anatomical model with Gaussian noise ($\sigma = 10$),

To investigate the influence of the parameters, we use the PSNR in decibels (dB) to quantify the quality of reconstructed images:

$$PSNR(\hat{\mathbf{x}}) = 10 \log_{10} \left(\frac{d^2}{MSE(\hat{\mathbf{x}})} \right), \quad MSE(\hat{\mathbf{x}}) = \frac{1}{|\Omega|} \sum_{\mathbf{v} \in \Omega} (\mathbf{x}(\mathbf{v}) - \hat{\mathbf{x}}(\mathbf{v}))^2 \quad (11)$$

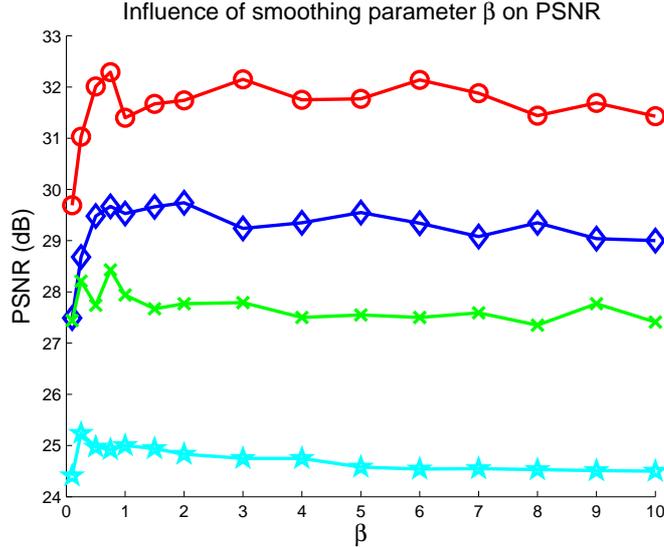


Figure 4: Influence of the smoothing parameter β on the PSNR computed with the reconstructed image compare the Brainweb ground truth. \circ : non-pathological noise free Brainweb image, \diamond : Multiple Sclerosis noise free Brainweb image, \times : non-pathological noisy Brainweb image, \star : Multiple Sclerosis noisy Brainweb image.

where MSE is the mean square error and d is the dynamic range of the image \mathbf{x} ($d = \max(\mathbf{x}) - \min(\mathbf{x}) = 255$).

In all experiments, a gradient descent method is used to optimize the cost function. The parameter λ is set to 0.01 (this value has been estimated in order to maximize the PSNR of the recovered image on Brainweb database).

By comparing the ground truth image with the reconstructed HR brain images, the Brainweb dataset allows us to evaluate the influence of each parameter of the proposed approach.

4.2 Influence of the smoothing parameter β

Figure 4 shows the influence of the smoothing parameter β for the computation of non-local weights w_{NLM} . For denoising purpose, Buades *et al.* [4] show that this parameter should be set to 1. Using the Brainweb image ground truth for normal and pathological cases, we investigate different values of β . Based on this experiment, it appears that a small value of β (such as 0.1 or 0.25) may significantly decrease the quality of the reconstructed image. However, for higher values of β , the quality of the reconstructed image does not seem to be strongly affected which leads to the assumption that the proposed non-local approach is not very sensitive to this setting.

4.3 Influence of the size of the search volume and the neighborhood

Figures 5 and 6 show the influence of the size of the search volume and the local neighborhood. Increasing the number of voxels in the search volume does not seem to affect the PSNR when the half-size of the search volume is greater than 3. Practically, using a small value for search volume size prevents useless computations. Concerning the patch size, it clearly appears that a patch of $3 \times 3 \times 3$ is of optimal size with respect to the PSNR and the computation time.

4.4 Influence of the image correlation

Figure 7 shows the correlation map computed between the HR T1-weighted image and the HR T2-weighted image for the multiple sclerosis Brainweb case. This figure shows that for regions containing outliers (i.e. MS lesions), α is low. For these regions, the final image is only reconstructed using information of LR T2-weighted image. Moreover, we also report results obtained with fixed values of α in Table 1 ($\alpha = 0$ or 1). When $\alpha = 0$ (which corresponds to the case of SR single-frame with non-local regularization), the quality of the reconstructed image is better than using interpolation methods. However, better results are obtained using an HR reference image during the reconstruction process (adaptive α using graph correlation or $\alpha = 1$).

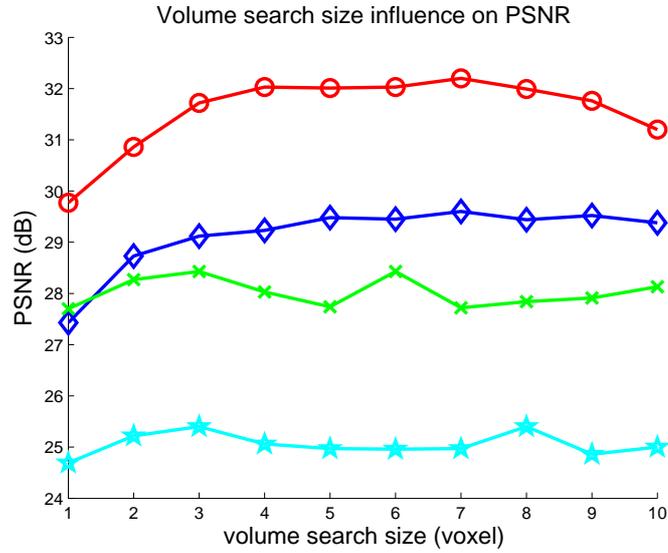


Figure 5: Influence of the size of the search volume on the PSNR computed with the reconstructed image compare the Brainweb ground truth. \circ : non-pathological noise free Brainweb image, \diamond : Multiple Sclerosis noise free Brainweb image, \times : non-pathological noisy Brainweb image, \star : Multiple Sclerosis noisy Brainweb image.

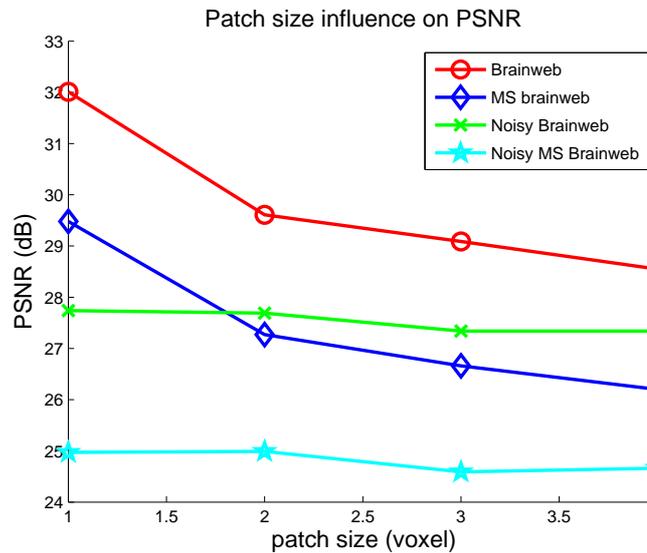


Figure 6: Influence of the size of the neighborhood on the PSNR computed with the reconstructed image compare the Brainweb ground truth. \circ : non-pathological noise free Brainweb image, \diamond : Multiple Sclerosis noise free Brainweb image, \times : non-pathological noisy Brainweb image, \star : Multiple Sclerosis noisy Brainweb image.

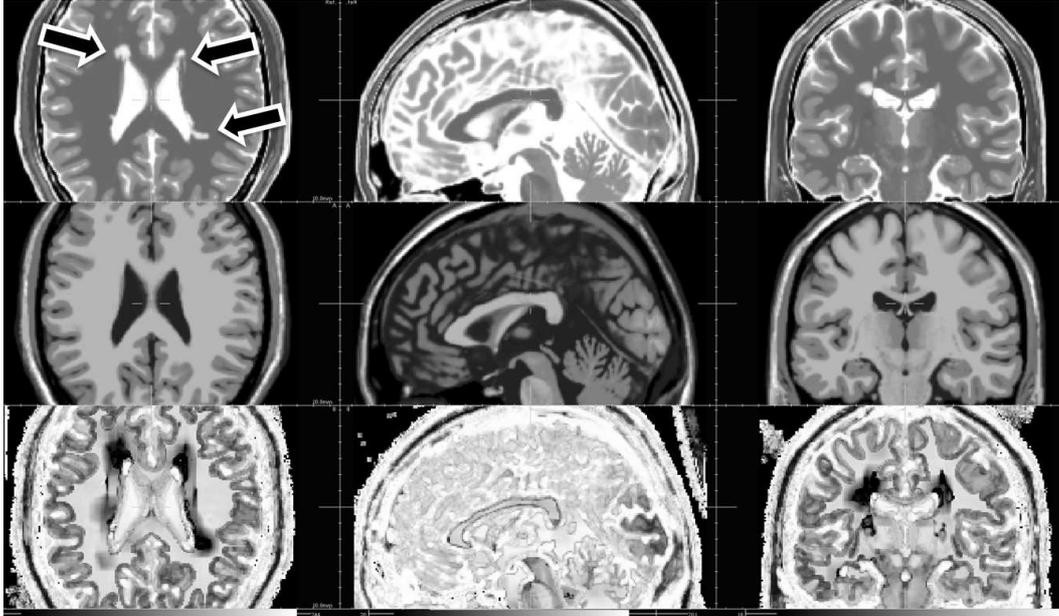


Figure 7: Estimated correlation map for the multiple sclerosis Brainweb image. First row : HR T2-weighted image, Second row: HR T1-weighted image, Third row: Correlation map (values of the parameter α . A linear intensity scale between 0 and 1 is used to display α). Three arrows show MS lesion locations (axial view).

4.5 Quantitative comparison with interpolation methods on Brainweb images

Table 1: Performances of image reconstruction methods (PSNR in dB) using noise free Brainweb images. The proposed method is compared with several interpolation methods. The influence of the correlation map (value of the parameter α) on the reconstruction results is also reported.

Method	Non pathological Image	Multiple Sclerosis Image
Nearest Neighbor Interpolation	23.74	20.89
Trilinear Interpolation	24.65	21.61
3rd order B-spline Interpolation	24.76	21.69
5th order B-spline Interpolation	24.73	21.67
Proposed method	32.20	29.48
Proposed method with $\alpha = 0$	28.30	26.05
Proposed method with $\alpha = 1$	32.09	29.16

We have reported PSNR results obtained with the different methods in Table 1. Results obtained with the proposed approach compare favorably with B-spline interpolation. Figures 8 and 9 show the results for non pathological and pathological MR Brainweb images for axial, coronal and sagittal views. Figure 10 is a zoom in on reconstruction results and allows a better visual comparison between interpolation method (trilinear and cubic B-spline) and the proposed SR method. It can be seen that details have been recovered and contrast between structures has been improved. Typical interpolation artefacts (see the skull for instance) are less visible on images reconstructed using the proposed method.

Figure 11 shows the difference between the T2-weighted ground truth image and the reconstructed images (non pathological case). It highlights the fact that the proposed framework provides a better reconstructed image, specifically at boundaries of brain structures. As pointed out in Section 3.3, the presence of brain lesions might introduce bias in reconstructed images using intermodality priors. Figure 12 shows results in MS lesion areas. Again, it appears that the proposed method using intermodality priors compares favorably with interpolation methods. More specifically, no spurious artefacts have been detected in reconstructed images. These results tend to show that our reconstruction method is robust to outliers such as MS lesions.

In MS, lesions are usually less visible in T1-weighted images than in T2-weighted images. To verify that our algorithm does not add any reconstruction artefacts, an HR T1-weighted image has been reconstructed using an HR T2-weighted image as reference image in a pathological case. Figure 12 shows details of reconstructed

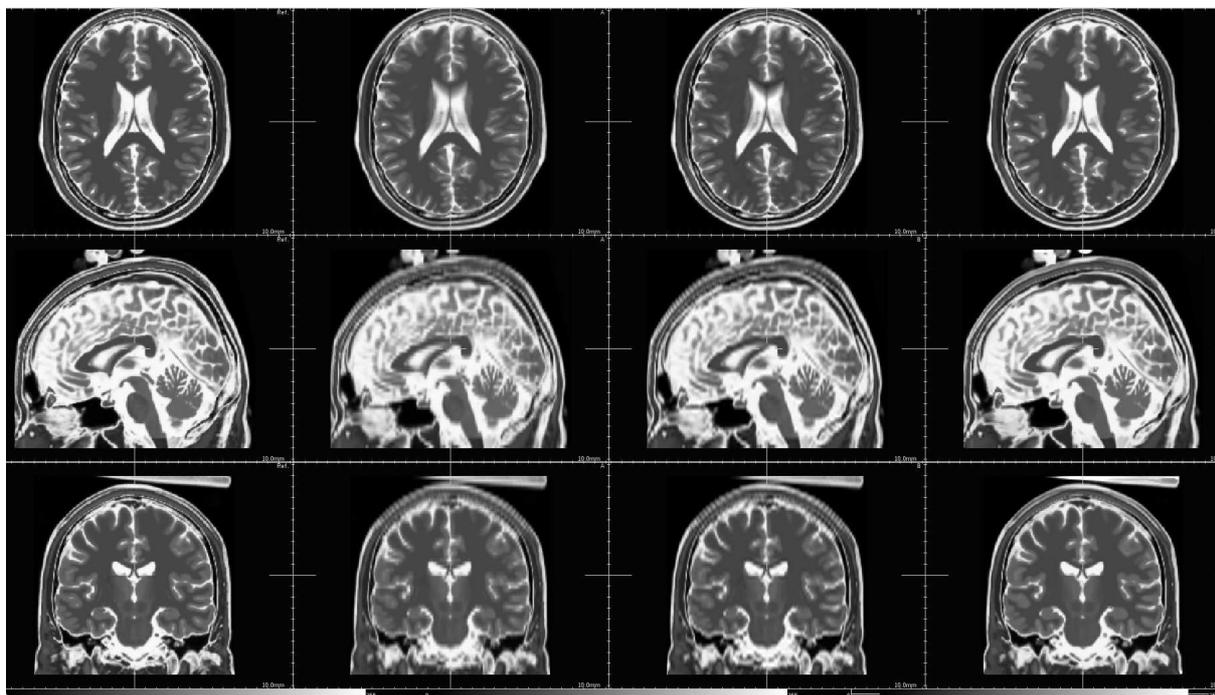


Figure 8: Reconstruction results (non pathological case). From left to right: Ground truth, Trilinear interpolation, third order B-spline interpolation, the proposed approach.

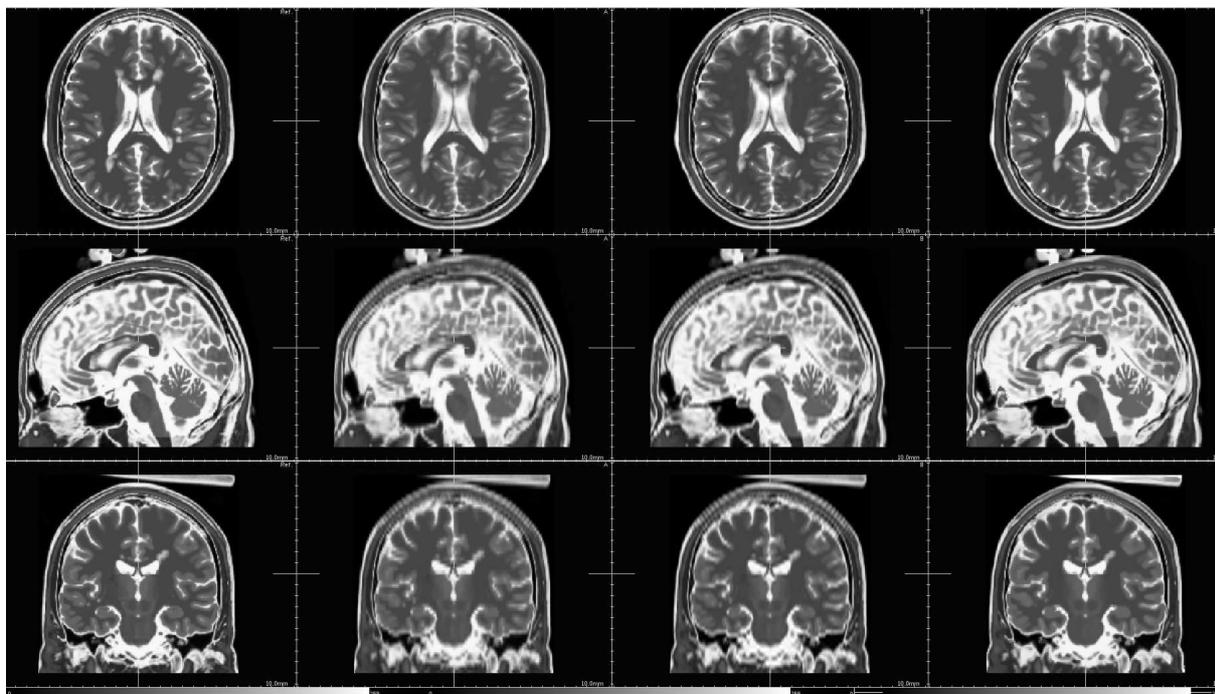


Figure 9: Reconstruction results in presence of MS lesions. From left to right: Ground truth, Trilinear interpolation, third order B-spline interpolation, the proposed approach.

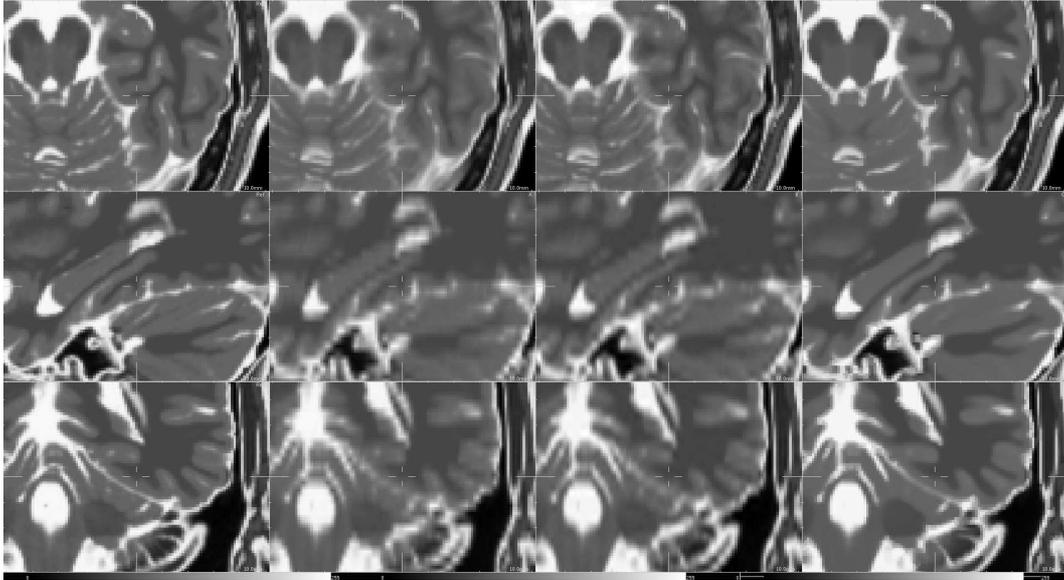


Figure 10: Zoom on reconstruction results. From left to right: Ground truth, Trilinear interpolation, third order B-spline interpolation, the proposed approach.

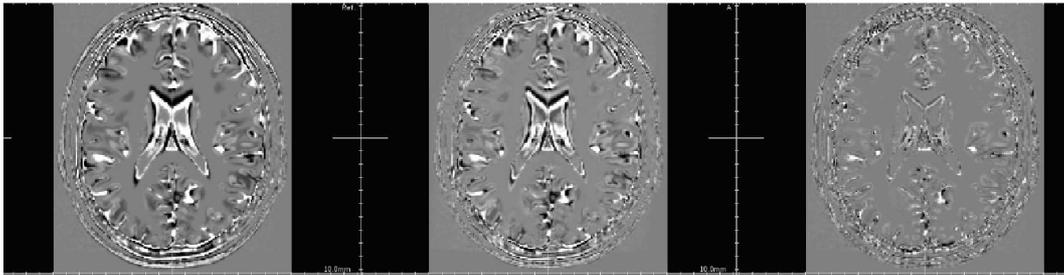


Figure 11: Difference between ground truth image and reconstruction results (non pathological case, axial view). From left to right: Trilinear interpolation, third order B-spline interpolation, the proposed approach. The same linear intensity scale is used for the display of image differences.

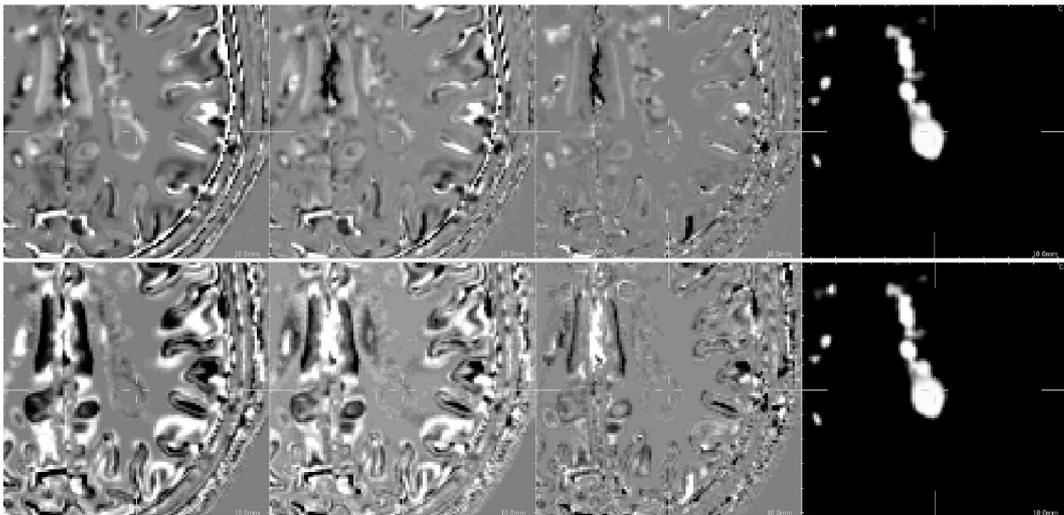


Figure 12: Difference between ground truth image and reconstruction results (pathological case). First row : reconstruction of a T2-weighted image, second row : reconstruction of a T1-weighted image. From left to right: Trilinear interpolation, third order B-spline interpolation, the proposed approach, MS lesion mask. The same linear intensity scale is used for the display of image differences.

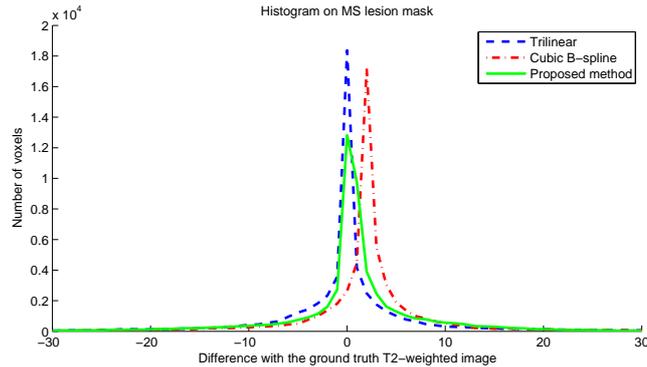


Figure 13: Histogram of error reconstruction (difference between ground truth T2-weighted image and reconstruction results) in MS lesion mask (pathological case). Mean and standard deviation of these distributions are: trilinear(0.32 ; 97.62), cubic B-spline(2.10 ; 71.17), proposed method(0.74 ; 67.72).

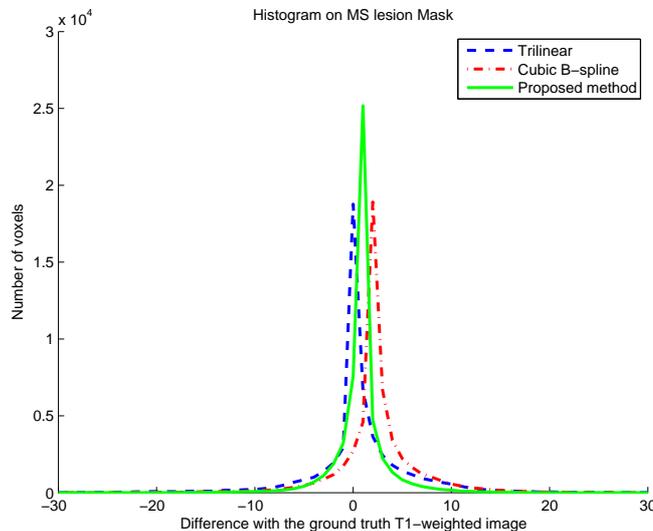


Figure 14: Histogram of error reconstruction (difference between ground truth T2-weighted image and reconstruction results) in MS lesion mask (pathological case). Mean and standard deviation of these distributions are: trilinear(0.64 ; 27.68), cubic B-spline(2.34 ; 18.10), proposed method(0.83 ; 7.71).

images in a MS lesion area. It seems that the proposed method does not add “hallucinated” (or false) lesion. This particular point is discussed in Section 5.

In addition to PSNR values and visual assessment, an error analysis has been done in MS lesion areas. Figures 13 and 14 show the histograms of error reconstruction for a reconstructed T2-weighted image (using an HR T1-weighted image as reference) and a reconstructed T1-weighted image (using an HR T2-weighted image as reference). In both cases, the standard deviation of the error reconstruction distribution is smaller for our method than for interpolation method. This tends to confirm that the proposed method allows a better reconstruction (specifically at boundaries) and that it is robust to outliers such as MS lesions.

4.6 Results on ADNI images

Data used in the preparation of this article were obtained from the Alzheimer’s Disease Neuroimaging Initiative (ADNI) database (www.loni.ucla.edu/ADNI). The ADNI was launched in 2003 by the National Institute on Aging (NIA), the National Institute of Biomedical Imaging and Bioengineering (NIBIB), the Food and Drug Administration (FDA), private pharmaceutical companies and non-profit organizations, as a \$60 million, 5-year public-private partnership. The primary goal of ADNI has been to test whether serial magnetic resonance imaging (MRI), positron emission tomography (PET), other biological markers, and clinical and neuropsychological assessment can be combined to measure the progression of mild cognitive impairment (MCI) and

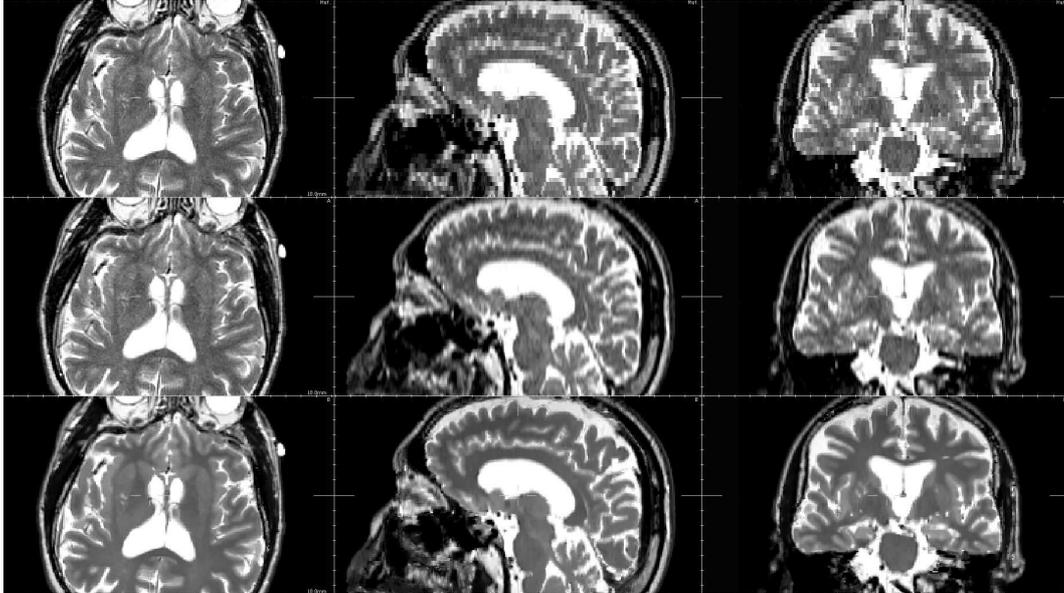


Figure 15: Reconstruction results on ADNI data (the LR T2-weighted image is firstly registered on the HR T1-weighted image). First row : nearest-neighbor interpolation, Second row : third order B-spline interpolation, Third row : the proposed approach.

early Alzheimer’s disease (AD). Determination of sensitive and specific markers of very early AD progression is intended to aid researchers and clinicians to develop new treatments and monitor their effectiveness, as well as lessen the time and cost of clinical trials. The Principle Investigator of this initiative is Michael W. Weiner, M.D., VA Medical Center and University of California – San Francisco. ADNI is the result of efforts of many co-investigators from a broad range of academic institutions and private corporations, and subjects have been recruited from over 50 sites across the U.S. and Canada. The initial goal of ADNI was to recruit 800 adults, ages 55 to 90, to participate in the research – approximately 200 cognitively normal older individuals to be followed for 3 years, 400 people with MCI to be followed for 3 years, and 200 people with early AD to be followed for 2 years.” For up-to-date information see www.adni-info.org.

The image size and resolution are respectively $160 \times 192 \times 192$ voxels and $1.2mm \times 1.250mm \times 1.250mm$ for the T1-weighted image, and $228 \times 256 \times 55$ voxels and $0.938mm \times 0.938mm \times 3.000mm$ for the LR axial T2-weighted image. The T1-weighted image is first registered to the LR T2-weighted image using a rigid transform and then the proposed SR technique is applied. Figures 15 and 16 show typical results obtained on the ADNI data using a T1-weighted image as the reference image to drive the reconstruction process of an axial T2-weighted image (the resolution of the reconstructed T2-weighted image is then : $0.938mm \times 0.938mm \times 0.938mm$). We can observe on these Figures that for the reconstructed image using the proposed approach, the contrast of brain structures is higher than using interpolation techniques. Typical interpolation artefacts (like staircase artefact) are less visible on the reconstructed image. Moreover, it can be noticed that the non-local regularization framework allows to jointly denoise and reconstruct the T2-weighted image.

To understand how the HR T1-weighted is used to drive the reconstruction process, the correlation map is displayed in Figure 17. It appears that the parameter α has a high value near to the edges which means that the HR reference image is mainly used for brain region with high contrast (interface between white matter and grey matter for instance). Conversely, uniform and/or textured regions such as caudate nucleus or putamen are only reconstructed using information existing in the LR T2-weighted image.

Looking at Figure 15, one can argue that the proposed reconstructed framework leads to an over smooth image which may not be realistic. As Figure 18 shows, the reconstructed T2-weighted image has indeed a very similar aspect to the HR T1-weighted image denoised using the Non-Local Mean algorithm [4]. Therefore, this property of the proposed algorithm appears to be beneficial since the proposed processing leads to a denoised HR T2-weighted MR image.

5 Discussion

The first contribution of this work concerns a simple idea for image SR which is the use of an HR reference image to improve the resolution of the LR image. As previously stated, such approach is directly related to

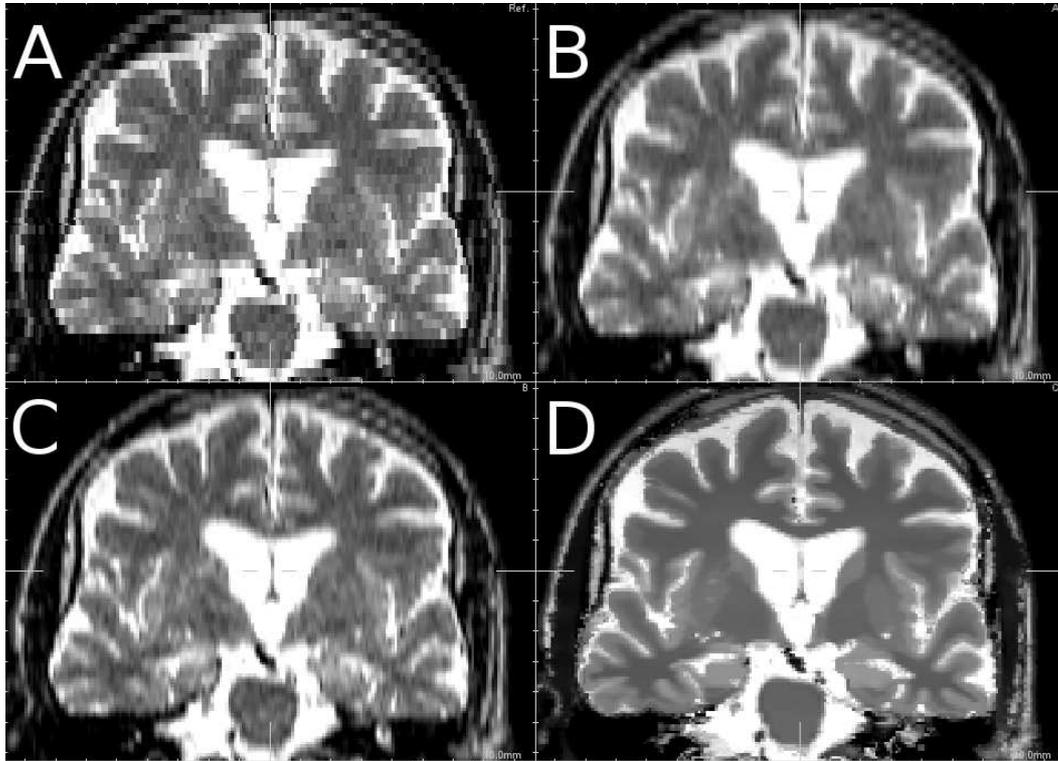


Figure 16: Zoom on reconstruction results on ADNI data (coronal view). A) Nearest Neighbor Interpolation, B) Trilinear Interpolation, C) third order B-spline interpolation, D) Proposed approach.

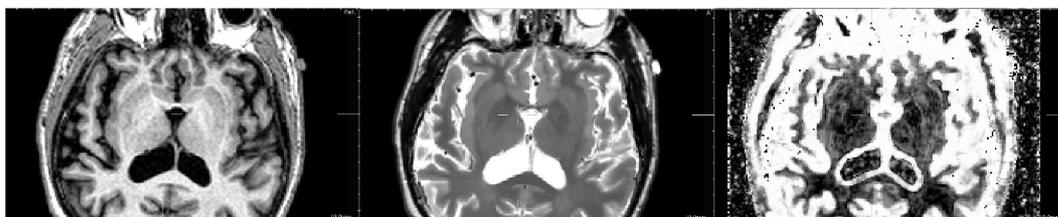


Figure 17: Estimated correlation map for ADNI images. First row : HR T1-weighted image, Second row: reconstructed T2-weighted image, Third row: Correlation map (values of the parameter α). A linear intensity scale between 0 and 1 is used to display α .

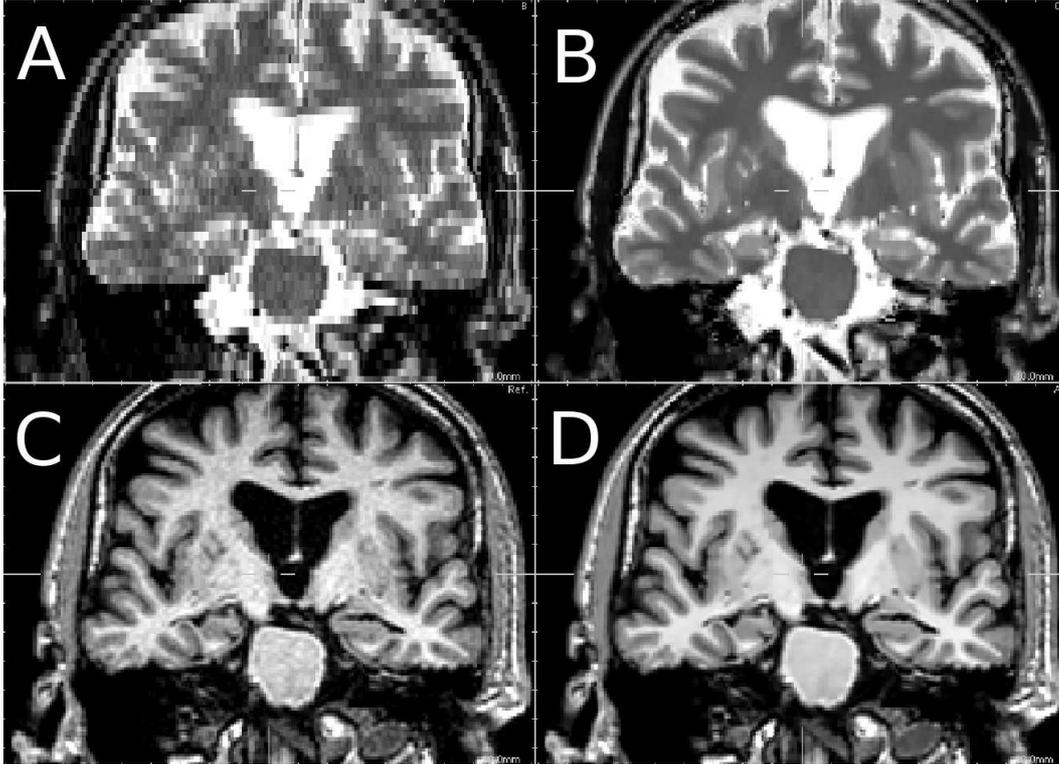


Figure 18: Zoom on reconstruction results on ADNI data (coronal view). A) Nearest Neighbor Interpolation, B) Proposed approach, C) T1-weighted image used as reference image, D) denoised T1-weighted image.

the medical context we are interested in. However, we believe that such new image SR approach may have a substantial impact in the image processing research field. Moreover, possible further work can focus on the unification of this framework with other previously proposed approaches such as object-based interpolation and registration-based interpolation.

The second contribution is about the modeling of the SR problem introducing a non-local regularization term. We have shown that example-based approaches such as non-local means can be embedded into the reconstruction process to enhance the performance of super resolution techniques. If an HR brain image of the patient is available, LR images of the same patient can be enhanced by exploiting non-local pairwise interactions. We believe that this work is a new original way to tackle the problem of image enhancement in MR imaging by exploiting the multimodality aspect of MR data. The key point of this work is the use of a non-local approach to define a new regularization term in the reconstruction process. The proposed methodology is related to example-based SR methods such as the one developed by Baker *et al.* [1] but in our case the learning database is reduced to one reference image. Moreover, in this work we exploit non-local interactions between voxels by defining non-local weights w (also called non-local graphs) which take into account non-local interactions in the reconstructed image and the reference image.

Experimental results show that the developed algorithm compares favorably with interpolation approaches. The two key points of the proposed approach, with respect to interpolation methods, are the use of an observation model (as in SR approaches) and the use of a reference HR image which drives the reconstruction process. Experiments on Brainweb images show that even with the presence of lesions visible only in the LR T2-weighted image, the reconstruction method we propose is able to recover such “outliers” without introducing artefacts which may come from the HR T1-weighted image (where lesions are not clearly visible). In this particular case, we would have thought that the lesions would disappear. Although our experiments tend to show that our approach is robust to such outliers, this is a crucial point that needs to be further investigated (small lesions, tumors, etc.). In general, this point raises the following question: what is the influence of an interpolation or image SR algorithm on image analysis (segmentation, detection, etc.) ?

This points out that such image enhancement technique needs to be validated using physical phantoms. Moreover, an evaluation on large database is required to prove that SR based methods can have a significant impact on medical image processing pipelines (specifically for registration, segmentation or change detection). Such validation can be application dependent and thus requires a specific evaluation framework. As it has been shown in Section 4, in addition to PSNR and visual assessment, there is a clear need to develop standard

criteria to measure image quality reconstruction of SR technique. This is a key point for image enhancement technique particularly in medical imaging where image artefacts have to be avoided.

On one hand, this work shows that SR based techniques provides higher MR image quality than standard interpolation algorithms. On the other hand, interpolation techniques do not require any rigid registration step and are less time consuming (SR based methods may require several hours per reconstruction). Based on our experiments (not shown in the paper), registration does not introduce artefacts in the reconstructed images using the proposed approach. If images are not correctly registered, the correlation coefficient α tends to zero which means that the reconstruction process is only driven by the T2-weighted image.

High resolution imaging is a key point for MR brain image analysis in order to study anatomical details. Since few brain image analysis algorithms take into account an observation model such as in Eq. 1, the proposed reconstruction approach may have a substantial impact on segmentation or registration. Therefore, such model based HR image reconstruction algorithm may represent an important step towards multimodal brain analysis at fine scale.

While in this work we are only focused on brain MR image reconstruction, the proposed approach relying intermodality priors might have potential applications to other medical image modalities. Future work would involve studying a similar method for multimodal images (computerized tomography, diffusion MRI, ultrasound etc.). Depending on the intended medical application, such approach could allow to save time of data acquisition by enabling image quality improvement as a post-acquisition step. The corner stone would concern then the criterion for linking the different modalities (How to calculate the local correlation map ? What type of information can be used between the imaging modalities ?).

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A Computation of weighted graphs $w_{\mathbf{x}}$ and $w_{\mathcal{E}_{\mathbf{x}}}$

$w_{\mathbf{x}}$ and $w_{\mathcal{E}_{\mathbf{x}}}$ are computed in the same way. For the sake of clarity, we describe only the computation of $w_{\mathbf{x}}$.

The weight $w_{\mathbf{x}}(\mathbf{v}, \mathbf{k})$ between two voxels \mathbf{v} and \mathbf{k} is computed as follows [7]:

$$w_{\mathbf{x}}(\mathbf{v}, \mathbf{k}) = \frac{1}{Z_{\mathbf{v}}} e^{-\frac{\|P(\mathbf{x}(\mathbf{v})) - P(\mathbf{x}(\mathbf{k}))\|_2^2}{2N_i\beta\sigma^2}} \quad (12)$$

where $P(\mathbf{x}(\mathbf{k}))$ is a 3D patch of \mathbf{x} centered in voxel \mathbf{k} and $P(\mathbf{x}(\mathbf{v}))$ is the 3D patch of \mathbf{x} centered in voxel \mathbf{v} ; $Z_{\mathbf{v}}$ is a constant of normalization; N_i is the number of voxels of a 3D patch. The distance between the 3D patches is the sum over voxels of patches of intensity differences using the L_2 norm. The specific case concerning the central point of the patch is computed as in [14]. With the assumption of Gaussian noise in images, β is set to 1 (see [4] for theoretical justifications) and the standard deviation of noise is estimated via pseudo-residuals $\epsilon_{\mathbf{v}}$ as defined in [10]. For each voxel \mathbf{v} of \mathbf{x} , let us define:

$$\epsilon_{\mathbf{v}} = \sqrt{\frac{6}{7}} \left(\mathbf{x}(\mathbf{v}) - \frac{1}{6} \sum_{\mathbf{k} \in \Omega(\mathbf{v})} \mathbf{x}(\mathbf{k}) \right) \quad (13)$$

where $\Omega(\mathbf{v})$ is the 6-neighborhood at voxel \mathbf{v} . The standard deviation of noise is computed as the least square estimator:

$$\hat{\sigma}^2 = \frac{1}{n} \sum_{\mathbf{v}} \epsilon_{\mathbf{v}}^2 \quad (14)$$

where n is the number of voxels of \mathbf{x} .

Moreover, as suggested by Mahmoudi and Sapiro in [16], voxel preselection can avoid useless computation and also improve the result of denoising. The preselection of relevant voxels is based on the mean and variance of patches [7]. $w_{\mathbf{x}}(\mathbf{v}, \mathbf{k})$ is set to 0 if one of these conditions is not fulfilled:

$$\mu < \frac{\overline{P(\mathbf{x}(\mathbf{v}))}}{\overline{P(\mathbf{x}(\mathbf{k}))}} < \frac{1}{\mu} \quad (15)$$

$$\sigma^2 < \frac{\text{var}(P(\mathbf{x}(\mathbf{v})))}{\text{var}(P(\mathbf{x}(\mathbf{k})))} < \frac{1}{\sigma^2} \quad (16)$$

with $\mu = 0.95$ and $\sigma^2 = 0.5$.

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